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p#16

RAW SEQUENCE LISTING

DATE: 10/30/2002

PATENT APPLICATION: US/09/673,490B

TIME: 16:02:48

Input Set : A:\13922.txt

Output Set: N:\CRF4\10302002\I673490B.raw

3 <110> APPLICANT: The University of Queensland
 5 <120> TITLE OF INVENTION: Novel omega conotoxin peptides
 7 <130> FILE REFERENCE: 2338740/MJC
 C--> 9 <140> CURRENT APPLICATION NUMBER: US/09/673,490B
 10 <141> CURRENT FILING DATE: 1999-04-16
 12 <150> PRIOR APPLICATION NUMBER: PCT/AU99/00288
 13 <151> PRIOR FILING DATE: 1999-04-16
 15 <160> NUMBER OF SEQ ID NOS: 44
 17 <170> SOFTWARE: PatentIn version 3.0
 19 <210> SEQ ID NO: 1
 20 <211> LENGTH: 6
 21 <212> TYPE: PRT
 22 <213> ORGANISM: conus catus
 24 <400> SEQUENCE: 1
 26 Ser Gly Thr Val Gly Arg
 27 1 5
 29 <210> SEQ ID NO: 2
 30 <211> LENGTH: 6
 31 <212> TYPE: PRT
 32 <213> ORGANISM: conus catus
 34 <400> SEQUENCE: 2
 36 Ser Lys Leu Met Tyr Asp
 37 1 5
 40 <210> SEQ ID NO: 3
 41 <211> LENGTH: 6
 42 <212> TYPE: PRT
 43 <213> ORGANISM: Artificial Sequence
 45 <220> FEATURE:
 46 <223> OTHER INFORMATION: Description of Artificial Sequence: A modified version of the
 second
 47 loop of CVID
 49 <400> SEQUENCE: 3
 51 Ser Arg Leu Met Tyr Asp
 52 1 5
 54 <210> SEQ ID NO: 4
 55 <211> LENGTH: 6
 56 <212> TYPE: PRT
 57 <213> ORGANISM: Artificial Sequence
 59 <220> FEATURE:
 60 <223> OTHER INFORMATION: Description of Artificial Sequence: A modified version of the
 second
 61 loop of CVID
 63 <400> SEQUENCE: 4
 65 Asp Arg Leu Met Tyr Asp

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66 1

5

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Input Set : A:\13922.txt

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68 <210> SEQ ID NO: 5
69 <211> LENGTH: 27
70 <212> TYPE: PRT
71 <213> ORGANISM: conus catus
73 <400> SEQUENCE: 5
75 Cys Lys Ser Lys Gly Ala Lys Cys Ser Lys Leu Met Tyr Asp Cys Cys
76 1          5          10          15
79 Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
80          20          25
83 <210> SEQ ID NO: 6
84 <211> LENGTH: 27
85 <212> TYPE: PRT
86 <213> ORGANISM: Artificial Sequence
88 <220> FEATURE:
89 <223> OTHER INFORMATION: Description of Artificial Sequence: A modified form of CVID
91 <400> SEQUENCE: 6
93 Cys Lys Ser Lys Gly Ala Lys Cys Ser Arg Leu Met Tyr Asp Cys Cys
94 1          5          10          15
96 Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
97          20          25
100 <210> SEQ ID NO: 7
101 <211> LENGTH: 27
102 <212> TYPE: PRT
103 <213> ORGANISM: Artificial Sequence
105 <220> FEATURE:
106 <223> OTHER INFORMATION: Description of Artificial Sequence: A modified form of CVID
108 <400> SEQUENCE: 7
110 Cys Lys Ser Lys Gly Ala Lys Cys Asp Arg Leu Met Tyr Asp Cys Cys
111 1          5          10          15
113 Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
114          20          25
116 <210> SEQ ID NO: 8
117 <211> LENGTH: 25
118 <212> TYPE: PRT
119 <213> ORGANISM: conus magus
121 <400> SEQUENCE: 8
123 Cys Lys Gly Lys Gly Ala Lys Cys Ser Arg Leu Met Tyr Asp Cys Cys
124 1          5          10          15
126 Thr Gly Ser Cys Arg Ser Gly Lys Cys
127          20          25
129 <210> SEQ ID NO: 9
130 <211> LENGTH: 26
131 <212> TYPE: PRT
132 <213> ORGANISM: conus magus
134 <400> SEQUENCE: 9
136 Cys Lys Gly Lys Gly Ala Pro Cys Arg Lys Thr Met Tyr Asp Cys Cys
137 1          5          10          15
139 Ser Gly Ser Cys Gly Arg Arg Gly Lys Cys
140          20          25

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142 <210> SEQ ID NO: 10
143 <211> LENGTH: 27
144 <212> TYPE: PRT
145 <213> ORGANISM: conus geographus
147 <220> FEATURE:
148 <221> NAME/KEY: misc_feature
149 <223> OTHER INFORMATION: Pro at positions 4, 10 and 21 is 4-Hyp
151 <400> SEQUENCE: 10
153 Cys Lys Ser Pro Gly Ser Ser Cys Ser Pro Thr Ser Tyr Asn Cys Cys
154 1 5 10 15
156 Arg Ser Cys Asn Pro Tyr Thr Lys Arg Cys Tyr
157 20 25
160 <210> SEQ ID NO: 11
161 <211> LENGTH: 18
162 <212> TYPE: DNA
163 <213> ORGANISM: conus catus
165 <400> SEQUENCE: 11
167 agcggcaccg taggtaga 18
170 <210> SEQ ID NO: 12
171 <211> LENGTH: 382
172 <212> TYPE: DNA
173 <213> ORGANISM: conus catus
175 <220> FEATURE:
176 <221> NAME/KEY: CDS
177 <222> LOCATION: (10)..(228)
179 <400> SEQUENCE: 12
181 atcatcaaaa atg aaa ctg acg tgt gtg gtg atc gtc gcc gtg ctg ctc ctg 51
182 Met Lys Leu Thr Cys Val Val Ile Val Ala Val Leu Leu Leu
183 1 5 10
185 acg gcc tgt caa ctc atc aca gct aat gac tcc aga ggt acg cag aag 99
186 Thr Ala Cys Gln Leu Ile Thr Ala Asn Asp Ser Arg Gly Thr Gln Lys
187 15 20 25 30
189 cat cgt gcc ctg agg tcg gac acc aaa ctc tcc atg tcg act cgc tgc 147
190 His Arg Ala Leu Arg Ser Asp Thr Lys Leu Ser Met Ser Thr Arg Cys
191 35 40 45
193 aag agt aaa gga gca aaa tgt tca aag ctt atg tat gac tgc tgc agc 195
194 Lys Ser Lys Gly Ala Lys Cys Ser Lys Leu Met Tyr Asp Cys Cys Ser
195 50 55 60
197 ggt tct tgc agc ggc acc gta ggt aga tgt ggc tgatccggcg cttgatctcc 248
198 Gly Ser Cys Ser Gly Thr Val Gly Arg Cys Gly
199 65 70
201 cccttctgtg ctctatcctt ttctgcctga gtccctcctta cctgagagtg gtcataaacc 308
203 actcatcacc tacccctgg aggtctcaaa gaactacttg aaataaagcc gcttgcaaaa 368
205 aaaaaaaaaa aaaa 382
208 <210> SEQ ID NO: 13
209 <211> LENGTH: 73
210 <212> TYPE: PRT
211 <213> ORGANISM: conus catus
213 <400> SEQUENCE: 13

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215 Met Lys Leu Thr Cys Val Val Ile Val Ala Val Leu Leu Leu Thr Ala
216 1 5 10 15
219 Cys Gln Leu Ile Thr Ala Asn Asp Ser Arg Gly Thr Gln Lys His Arg
220 20 25 30
223 Ala Leu Arg Ser Asp Thr Lys Leu Ser Met Ser Thr Arg Cys Lys Ser
224 35 40 45
227 Lys Gly Ala Lys Cys Ser Lys Leu Met Tyr Asp Cys Cys Ser Gly Ser
228 50 55 60
231 Cys Ser Gly Thr Val Gly Arg Cys Gly
232 65 70
235 <210> SEQ ID NO: 14
236 <211> LENGTH: 27
237 <212> TYPE: PRT
238 <213> ORGANISM: conus catus
240 <400> SEQUENCE: 14
242 Cys Arg Ser Lys Gly Ala Lys Cys Ser Lys Leu Met Tyr Asp Cys Cys
243 1 5 10 15
245 Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
246 20 25
248 <210> SEQ ID NO: 15
249 <211> LENGTH: 27
250 <212> TYPE: PRT
251 <213> ORGANISM: conus catus
253 <400> SEQUENCE: 15
255 Cys Lys Ser Lys Gly Ala Arg Cys Ser Lys Leu Met Tyr Asp Cys Cys
256 1 5 10 15
258 Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
259 20 25
261 <210> SEQ ID NO: 16
262 <211> LENGTH: 27
263 <212> TYPE: PRT
264 <213> ORGANISM: conus catus
266 <400> SEQUENCE: 16
268 Cys Lys Ser Lys Gly Ala Gln Cys Ser Lys Leu Met Tyr Asp Cys Cys
269 1 5 10 15
271 Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
272 20 25
274 <210> SEQ ID NO: 17
275 <211> LENGTH: 27
276 <212> TYPE: PRT
277 <213> ORGANISM: conus catus
279 <400> SEQUENCE: 17
281 Cys Lys Ser Lys Gly Ala Lys Cys Ser Lys Leu Met Tyr Asp Cys Cys
282 1 5 10 15
285 Ser Gly Ser Cys Ser Gly Ala Val Gly Arg Cys
286 20 25
288 <210> SEQ ID NO: 18
289 <211> LENGTH: 27
290 <212> TYPE: PRT

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RAW SEQUENCE LISTING

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291 <213> ORGANISM: Artificial Sequence
293 <220> FEATURE:
294 <223> OTHER INFORMATION: Description of Artificial Sequence: A derivative of CVID
296 <400> SEQUENCE: 18
298 Cys Lys Ser Lys Gly Ala Lys Cys Asp Lys Leu Met Tyr Asp Cys Cys
299 1      5      10      15
301 Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
302      20      25
304 <210> SEQ ID NO: 19
305 <211> LENGTH: 27
306 <212> TYPE: PRT
307 <213> ORGANISM: Artificial Sequence
309 <220> FEATURE:
310 <223> OTHER INFORMATION: Description of Artificial Sequence: A derivative of CVID
312 <400> SEQUENCE: 19
314 Cys Lys Tyr Lys Gly Ala Lys Cys Ser Arg Leu Met Tyr Asp Cys Cys
315 1      5      10      15
317 Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
318      20      25
320 <210> SEQ ID NO: 20
321 <211> LENGTH: 27
322 <212> TYPE: PRT
323 <213> ORGANISM: Artificial Sequence
325 <220> FEATURE:
326 <223> OTHER INFORMATION: Description of Artificial Sequence: A derivative of CVID
328 <400> SEQUENCE: 20
330 Cys Lys Ser Lys Gly Ala Lys Cys Ser Lys Leu Ala Tyr Asp Cys Cys
331 1      5      10      15
333 Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
334      20      25
336 <210> SEQ ID NO: 21
337 <211> LENGTH: 27
338 <212> TYPE: PRT
339 <213> ORGANISM: Artificial Sequence
341 <220> FEATURE:
342 <223> OTHER INFORMATION: Description of Artificial Sequence: A derivative of CVID
344 <400> SEQUENCE: 21
346 Cys Lys Ser Lys Gly Ala Lys Cys Ser Lys Leu Met Tyr Asp Cys Cys
347 1      5      10      15
349 Thr Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
350      20      25
352 <210> SEQ ID NO: 22
353 <211> LENGTH: 27
354 <212> TYPE: PRT
355 <213> ORGANISM: Artificial Sequence
357 <220> FEATURE:
358 <223> OTHER INFORMATION: Description of Artificial Sequence: A derivative of CVID
360 <220> FEATURE:
361 <221> NAME/KEY: misc_feature

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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 10/30/2002
PATENT APPLICATION: US/09/673,490B TIME: 16:02:49

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:22; Xaa Pos. 5
Seq#:30; Xaa Pos. 12
Seq#:31; Xaa Pos. 12
Seq#:37; Xaa Pos. 4
Seq#:38; Xaa Pos. 4
Seq#:39; Xaa Pos. 2,3,4,6,7,9,12,14,17,22

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:33; Line(s) 567
Seq#:34; Line(s) 581
Seq#:35; Line(s) 595
Seq#:36; Line(s) 613
Seq#:37; Line(s) 631
Seq#:38; Line(s) 650